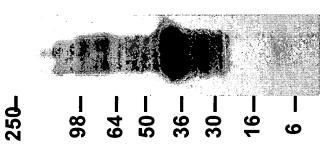


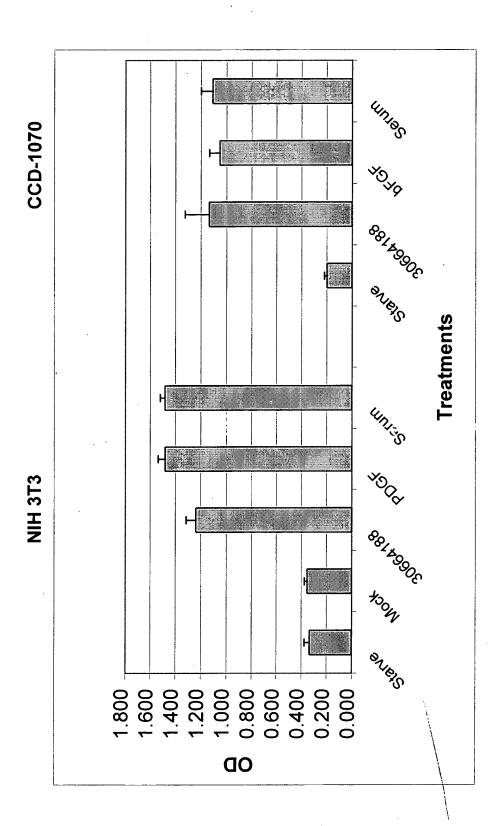
FIG. 1.

Multiple Alignment:

8 7 8	119 114	179 165	238 214	296 274	356 331	330 345
I WHELFFYYTLICANFCSCRDTSATPOSASIKALRNANLRRDESNHLTDLYRRDETTOWKG	61 NGYWOSPRFPNSWPRNLÜLTWRLHS-QENTRIQLVFDNOFGLEEAENDICRYDFYEVEDE	SETSTIBRERWCGHKEVPPRIKSRTNQIKITEKSDDVFVAKPGFKIYVSLEDFQPAAAS	180 ETNWESVTSSTSGVSYNSPSVTDP-TLIADALDKKIAEFDTVEDLEKYFNP ESWOEDLEN) MWLDTPRYRGRSYHD. RKS. KVDIDRINDDAKRKSCTPRNKSVNIREELKLANVFFFRC	297 LLV QRCGGNCGCGTVNWRSCTCNSGKTVKKYHEVLQFEPGHIKRRGRAKTMALVDNQLDH	HERCDONCRSSRPPR HEECDOVCRSTGG
IMSLFGLLLLTSALAGOROGROGROSNLSSKFOFSSNKEONGRODPO-HERIITWST	55 NGS THSPRFPHMYPRNTWLVWRLYAWEENVWIQLTFDERFGLEDPEDDICKYDFVEVEEP	SDG TILBRWCGSGTVPGKQISKGNQIRIRFVSDDVFPSEPGFCIHVNLVMP	166QFTEAVSPSVLPPSALPLDLINNAITAFSTLEDLIRYLEPERWOLDLED	5 LKRPTWQLLGKARVFBRKSRVWDINLITEEVRLKSCTPRNFSVSIREELKRTDTIFWFGC	275 LLV KRCGGNCACCLHNCNECQCVPSKVTKKYHEVLQLRPKTGVRGLHK SLTDYALEH	
		<u>1</u>	⊠ ÿ	23		337
30664188.0.99	30664188.0.99	30664188.0.99 120	30664188.0.99	30664188.0.99 239	30664188.0.99	30664188.0.99
VEGF-E	VEGF-E	VEGF-E 115	VEGF-E	VEGF-E 215	VEGF-E	VEGF-E



Imidazole Elution



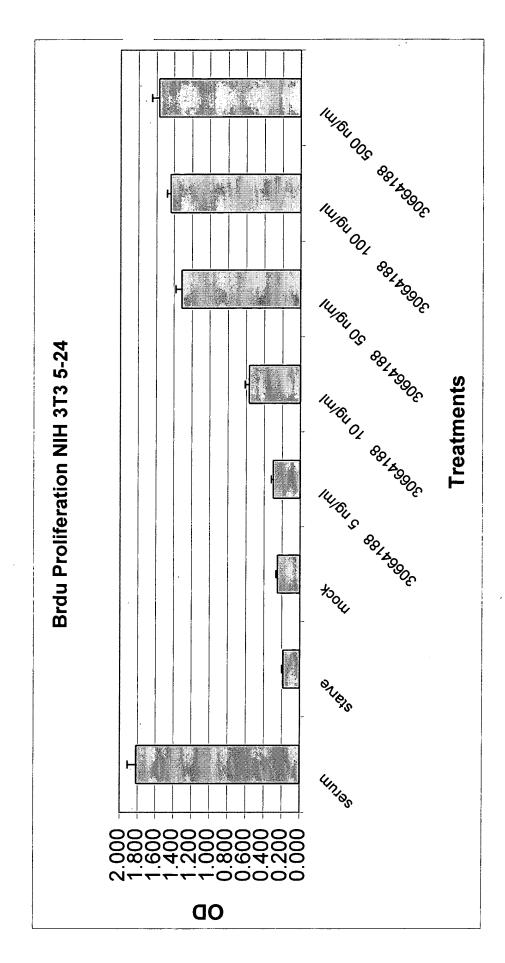
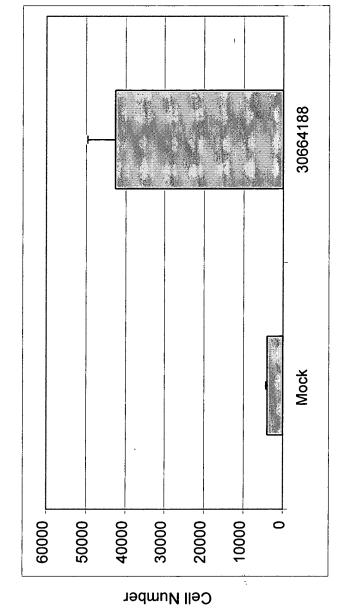
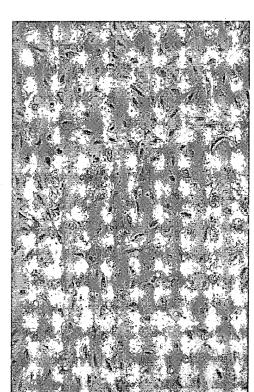


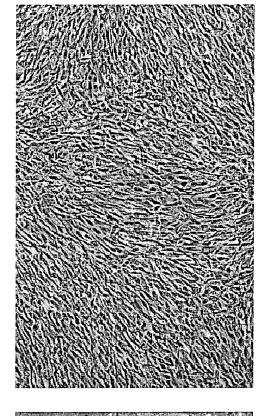
FIG. 7.



pCEP4sec CM

pCEP4sec(30664188) CM





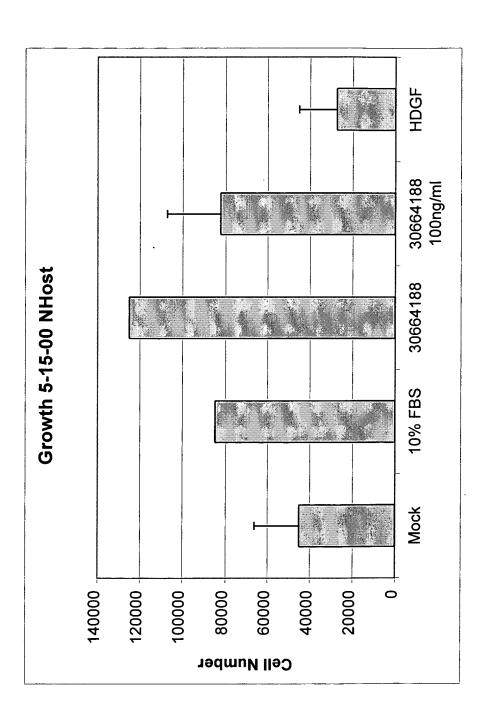
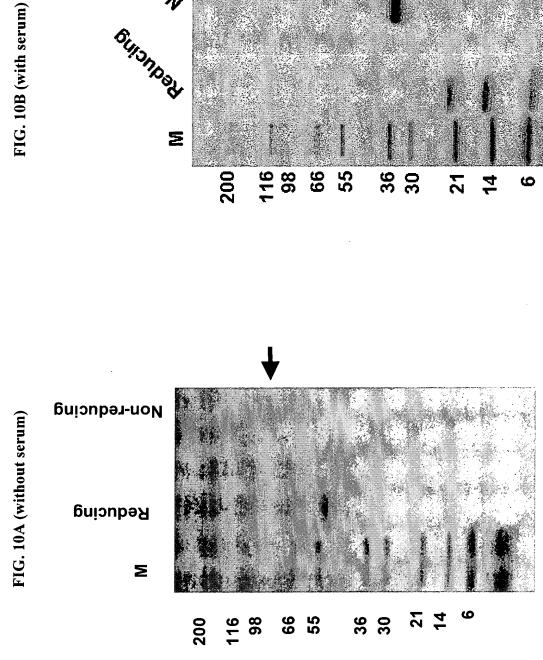
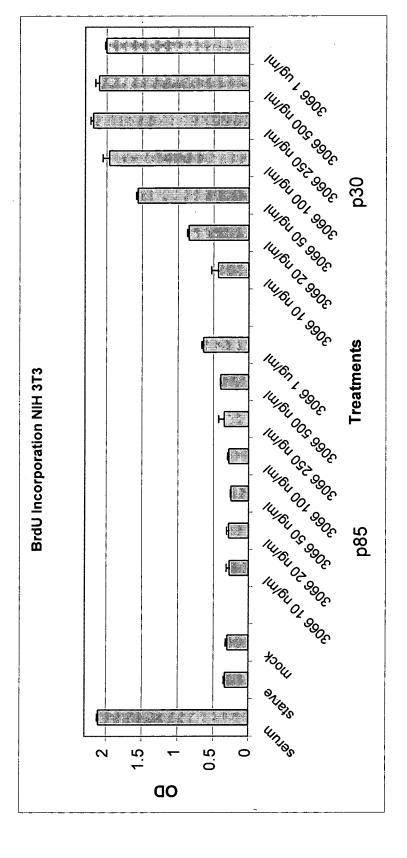


FIG. 10.





10 / 11